# RAW SEQUENCE LISTING
# ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _09/938,703A_

Source: _1600,_

Date Processed by STIC: _10/29/2003_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
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2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
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     http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/2003):
    U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/2003

1600

RAW SEQUENCE LISTING                     DATE: 10/29/2003
PATENT APPLICATION: **US/09/938,703A**      TIME: 15:11:12

Input Set : **A:\2023d.ST25.txt**
Output Set: **N:\CRF4\10292003\I938703A.raw**

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 3 <110> APPLICANT: Samson, Michael
 4        Parmentier, Marc
 5        Vassart, Gilbert
 6        Libert, Frederick
 8 <120> TITLE OF INVENTION: Therapeutic Applications for CCR Chemokine Receptors
10 <130> FILE REFERENCE: 9409/2023D
```
C--> `12 <140> CURRENT APPLICATION NUMBER: US/09/938,703A`
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13 <141> CURRENT FILING DATE: 2001-08-24
15 <150> PRIOR APPLICATION NUMBER: 09/626,939
16 <151> PRIOR FILING DATE: 2000-07-27
18 <150> PRIOR APPLICATION NUMBER: 09/08/833,752
19 <151> PRIOR FILING DATE: 1997-04-09
21 <150> PRIOR APPLICATION NUMBER: EP 96870021.1
22 <151> PRIOR FILING DATE: 1996-03-01
24 <150> PRIOR APPLICATION NUMBER: EP 96870102.9
25 <151> PRIOR FILING DATE: 1996-08-06
27 <160> NUMBER OF SEQ ID NOS: 18
29 <170> SOFTWARE: PatentIn version 3.1
31 <210> SEQ ID NO: 1
32 <211> LENGTH: 792
33 <212> TYPE: DNA
34 <213> ORGANISM: Homo sapiens
36 <400> SEQUENCE: 1
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39 tcccttcact acaaaacttc attgcttggc caaaaagaga gttaattcaa tgtagacatc   120
41 tatgtaggca attaaaaacc tattgatgta taaaacagtt tgcattcatg gagggcaact   180
43 aaatacattc taggacttta taaaagatca ctttttattt atgcacaggg tggaacaaga   240
45 tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc   300
47 aaaaaatcaa tgtgaagcaa atcgcagccc gcctcctgcc tccgctctac tcactggtgt   360
49 tcatctttgg ttttgtgggc aacatgctgg tcatcctcat cctgataaac tgcaaaaggc   420
51 tgaagagcat gactgacatc tacctgctca acctggccat ctctgacctg ttttttccttc   480
53 ttactgtccc cttctgggct cactatgctg ccgcccagtg ggactttgga aatacaatgt   540
55 gtcaactctt gacagggctc tatttatag gcttcttctc tggaatcttc ttcatcatcc    600
57 tcctgacaat cgataggtac ctggctgtcg tccatgctgt gtttgcttta aaagccagga   660
59 cggtcacctt tggggtggtg acaagtgtga tcacttgggt ggtggctgtg tttgcgtctc   720
61 tcccaggaat catctttacc agatctcaaa aagaaggtct tcattacacc tgcagctctc   780
63 attttccata ca                                                       792
66 <210> SEQ ID NO: 2
67 <211> LENGTH: 1477
68 <212> TYPE: DNA
69 <213> ORGANISM: Homo sapiens
71 <220> FEATURE:
72 <221> NAME/KEY: misc_feature
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Does Not Comply
Corrected Diskette Needed

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PATENT APPLICATION: **US/09/938,703A**      TIME: 15:11:12

Input Set : **A:\2023d.ST25.txt**
Output Set: **N:\CRF4\10292003\I938703A.raw**

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 74 <223> OTHER INFORMATION: Any nucleotide
 77 <220> FEATURE:
 78 <221> NAME/KEY: misc_feature
 79 <222> LOCATION: (1384)..(1385)
 80 <223> OTHER INFORMATION: Any nucleotide
 83 <400> SEQUENCE: 2
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 86 tcccttcact acaaaacttc attgcttggc caaaaagaga gttaattcaa tgtagacatc     120
 88 tatgtaggca attaaaaacc tattgatgta taaaacagtt tgcattcatg gagggcaact     180
 90 aaatacattc taggacttta taaaagatca ctttttattt atgcacaggg tggaacaaga     240
 92 tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc     300
 94 aaaaaatcaa tgtgaagcaa atcgcagccc gcctcctgcc tccgctctac tcactggtgt     360
 96 tcatctttgg ttttgtgggc aacatgctgg tcatcctcat cctgataaac tgcaaaaggc     420
 98 tgaagagcat gactgacatc tacctgctca acctggccat ctctgacctg tttttccttc     480
100 ttactgtccc cttctgggct cactatgctg ccgcccagtg ggactttgga aatacaatgt     540
102 gtcaactctt gacagggctc tattttatag gcttcttctc tggaatcttc ttcatcatcc     600
104 tcctgacaat cgataggtac ctggctgtcg tccatgctgt gtttgcttta aaagccagga     660
106 cggtcacctt tggggtggtg acaagtgtga tcacttgggt ggtggctgtg tttgcgtctc     720
108 tcccaggaat catctttacc agatctcaaa aagaaggtct tcattacacc tgcagctctc     780
110 attttccata cagtcagtat caattctgga agaatttcca gacattaaag atagtcatct     840
112 tggggctggt cctgccgctg cttgtcatgg tcatctgcta ctcgggaatc ctaaaaactc     900
114 tgcttcggtg tcgaaatgag aagaagaggc acagggctgt gaggcttatc ttcaccatca     960
116 tgattgttta ttttctcttc tgggctccct acaacattgt ccttctcctg aacaccttcc    1020
118 aggaattctt tggcctgaat aattgcagta gctctaacag gttggaccaa gctatgcagg    1080
120 tgacagagac tcttgggatg acgcactgct gcatcaaccc catcatctat gcctttgtcg    1140
122 gggagaagtt cagaaactac ctcttagtct tcttccaaaa gcacattgcc aaacgcttct    1200
124 gcaaatgctg ttctattttc cagcaagagg ctcccgagcg agcaagctca gtttacaccc    1260
126 gatccactgg ggagcaggaa atatctgtgg gcttgtgaca cggactcaag tgggctggtg    1320
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**W--> 128 acccagtcag agttgtgcac atggcttagt tttcatacac agcctgggct ggggggtnggt 1380**

```
130 tggnngaggt cttttttaaa aggaagttac tgttatagag ggtctaagat tcatccattt    1440
132 atttggcatc tgtttaaagt agattagatc cgaattc                             1477
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136 <211> LENGTH: 1442
137 <212> TYPE: DNA
138 <213> ORGANISM: Homo sapiens
140 <400> SEQUENCE: 3
141 gaattccccc aacagagcca agctctccat ctagtggaca gggaagctag cagcaaacct      60
143 tcccttcact acaaaacttc attgcttggc caaaaagaga gttaattcaa tgtagacatc     120
145 tatgtaggca attaaaaacc tattgatgta taaaacagtt tgcattcatg gagggcaact     180
147 aaatacattc taggacttta taaaagatca ctttttattt atgcacaggg tggaacaaga     240
149 tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc     300
151 aaaaaatcaa tgtgaagcaa atcgcagccc gcctcctgcc tccgctctac tcactggtgt     360
153 tcatctttgg ttttgtgggc aacatgctgg tcatcctcat cctgataaac tgcaaaaggc     420
155 tgaagagcat gactgacatc tacctgctca acctggccat ctctgacctg tttttccttc     480
157 ttactgtccc cttctgggct cactatgctg ccgcccagtg ggactttgga aatacaatgt     540
159 gtcaactctt gacagggctc tattttatag gcttcttctc tggaatcttc ttcatcatcc     600
161 tcctgacaat cgataggtac ctggctgtcg tccatgctgt gtttgcttta aaagccagga     660
163 cggtcacctt tggggtggtg acaagtgtga tcacttgggt ggtggctgtg tttgcgtctc     720
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RAW SEQUENCE LISTING                    DATE: 10/29/2003
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Input Set : **A:\2023d.ST25.txt**
Output Set: **N:\CRF4\10292003\I938703A.raw**

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165 tcccaggaat catctttacc agatctcaaa aagaaggtct tcattacacc tgcagctctc   780
167 attttccata cattaaagat agtcatcttg gggctggtcc tgccgctgct tgtcatggtc   840
169 atctgctact cgggaatcct aaaaactctg cttcggtgtc gaaatgagaa gaagaggcac   900
171 agggctgtga ggcttatctt caccatcatg attgtttatt ttctcttctg ggctccctac   960
173 aacattgtcc ttctcctgaa caccttccag gaattctttg gcctgaataa ttgcagtagc  1020
175 tctaacaggt tggaccaagc tatgcaggtg acagagactc ttgggatgac gcactgctgc  1080
177 atcaaccсca tcatctatgc ctttgtcggg gagaagttca gaaactacct cttagtcttc  1140
179 ttccaaaagc acattgccaa acgcttctgc aaatgctgtt ctattttcca gcaagaggct  1200
181 cccgagcgag caagctcagt ttacacccga tccactgggg agcaggaaat atctgtgggc  1260
183 ttgtgacacg gactcaagtg ggctggtgac ccagtcagag ttgtgcacat ggcttagttt  1320
185 tcatacacag cctgggctgg gggtggttgg gaggtctttt ttaaaaggaa gttactgtta  1380
187 tagagggtct aagattcatc catttatttg gcatctgttt aaagtagatt agatccgaat  1440
189 tc                                                                 1442
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193 <211> LENGTH: 184
194 <212> TYPE: PRT
195 <213> ORGANISM: Homo sapiens
197 <400> SEQUENCE: 4
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203 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
204             20                  25                  30
207 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
208         35                  40                  45
211 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
212     50                  55                  60
215 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
216 65                  70                  75                  80
219 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
220                 85                  90                  95
223 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
224             100                 105                 110
227 Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu
228         115                 120                 125
231 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
232     130                 135                 140
235 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
236 145                 150                 155                 160
239 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
240             165                 170                 175
243 Thr Cys Ser Ser His Phe Pro Tyr
244             180
247 <210> SEQ ID NO: 5
248 <211> LENGTH: 352
249 <212> TYPE: PRT
250 <213> ORGANISM: Homo sapiens
252 <400> SEQUENCE: 5
254 Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
255 1               5                   10                  15
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258 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
259          20                  25                  30
262 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
263          35                  40                  45
266 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
267      50                  55                  60
270 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
271 65                  70                  75                  80
274 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
275              85                  90                  95
278 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
279          100                 105                 110
282 Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu
283          115                 120                 125
286 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
287      130                 135                 140
290 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
291 145                 150                 155                 160
294 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
295              165                 170                 175
298 Thr Cys Ser Ser His Phe Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn
299          180                 185                 190
302 Phe Gln Thr Leu Lys Ile Val Ile Leu Gly Leu Val Leu Pro Leu Leu
303          195                 200                 205
306 Val Met Val Ile Cys Tyr Ser Gly Ile Leu Lys Thr Leu Leu Arg Cys
307      210                 215                 220
310 Arg Asn Glu Lys Lys Arg His Arg Ala Val Arg Leu Ile Phe Thr Ile
311 225                 230                 235                 240
314 Met Ile Val Tyr Phe Leu Phe Trp Ala Pro Tyr Asn Ile Val Leu Leu
315              245                 250                 255
318 Leu Asn Thr Phe Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser Ser
319          260                 265                 270
322 Asn Arg Leu Asp Gln Ala Met Gln Val Thr Glu Thr Leu Gly Met Thr
323          275                 280                 285
326 His Cys Cys Ile Asn Pro Ile Ile Tyr Ala Phe Val Gly Glu Lys Phe
327      290                 295                 300
330 Arg Asn Tyr Leu Leu Val Phe Phe Gln Lys His Ile Ala Lys Arg Phe
331 305                 310                 315                 320
334 Cys Lys Cys Cys Ser Ile Phe Gln Gln Glu Ala Pro Glu Arg Ala Ser
335              325                 330                 335
338 Ser Val Tyr Thr Arg Ser Thr Gly Glu Gln Glu Ile Ser Val Gly Leu
339          340                 345                 350
342 <210> SEQ ID NO: 6
343 <211> LENGTH: 215
344 <212> TYPE: PRT
345 <213> ORGANISM: Homo sapiens
347 <400> SEQUENCE: 6
349 Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
350 1               5                   10                  15
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PATENT APPLICATION:  US/09/938,703A      TIME: 15:11:12

Input Set : A:\2023d.ST25.txt
Output Set: N:\CRF4\10292003\I938703A.raw

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353 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
354             20                  25                  30
357 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
358         35                  40                  45
361 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
362     50                  55                  60
365 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
366 65                  70                  75                  80
369 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
370             85                  90                  95
373 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
374             100                 105                 110
377 Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu
378         115                 120                 125
381 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
382     130                 135                 140
385 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
386 145                 150                 155                 160
389 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
390             165                 170                 175
393 Thr Cys Ser Ser His Phe Pro Tyr Ile Lys Asp Ser His Leu Gly Ala
394             180                 185                 190
397 Gly Pro Ala Ala Ala Cys His Gly His Leu Leu Leu Gly Asn Pro Lys
398         195                 200                 205
401 Asn Ser Ala Ser Val Ser Lys
402     210                 215
405 <210> SEQ ID NO: 7
406 <211> LENGTH: 360
407 <212> TYPE: PRT
408 <213> ORGANISM: Homo sapiens
410 <220> FEATURE:
411 <221> NAME/KEY: MISC_FEATURE
412 <222> LOCATION: (325)..(327)
413 <223> OTHER INFORMATION: Xaa = any amino acid
416 <400> SEQUENCE: 7
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419 1               5                   10                  15
422 Gly Glu Glu Val Thr Thr Phe Phe Asp Tyr Asp Tyr Gly Ala Pro Cys
423             20                  25                  30
426 His Lys Phe Asp Val Lys Gln Ile Gly Ala Gln Leu Leu Pro Pro Leu
427         35                  40                  45
430 Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn Met Leu Val Val
431     50                  55                  60
434 Leu Ile Leu Ile Asn Cys Lys Lys Leu Lys Cys Leu Thr Asp Ile Tyr
435 65                  70                  75                  80
438 Leu Leu Asn Leu Ala Ile Ser Asp Leu Leu Phe Ile Ile Thr Leu Pro
439             85                  90                  95
442 Leu Trp Ala His Ser Ala Ala Asn Glu Trp Val Phe Gly Asn Ala Met
443             100                 105                 110
```

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<212> TYPE: DNA
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<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: :
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<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: :
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<210> SEQ ID NO 17
<211> LENGTH: 27
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: :
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     ctgatctaga gccatgtgca caactct                                    27
```

*FYI*

Input Set : **A:\2023d.ST25.txt**
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**Please Note:**
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; N Pos. 1377,1384,1385
Seq#:7; Xaa Pos. 325,326,327
Seq#:8; Xaa Pos. 231,232,233,333,334,335
Seq#:10; Xaa Pos. 145,146,147,321,322,323

**Use of <220> Feature(NEW RULES):**   *error explanation*
**Sequence(s)__are missing the <220> Feature and associated headings.**
**Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence"**
**or"Unknown". Please explain source of genetic material in <220> to <223>**
**section (See "Federal Register," 6/01/98, Vol. 63, No. 104,pp.29631-32)**
**(Sec.1.823 of new Rules)**

Seq#:14,15,16,17

Input Set : **A:\2023d.ST25.txt**

Output Set: **N:\CRF4\10292003\I938703A.raw**

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L:12 M:270 C: Current Application Number differs, Replaced Current Application Number
L:128 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1320
M:341 Repeated in SeqNo=2
L:498 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:320
L:585 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:224
M:341 Repeated in SeqNo=8
L:775 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:144
M:341 Repeated in SeqNo=10
L:889 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:891 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:14, <213>
ORGANISM:Artificial Sequence
L:891 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:14, <213>
ORGANISM:Artificial Sequence
L:891 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:14,Line#:891
L:898 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:900 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:15, <213>
ORGANISM:Artificial Sequence
L:900 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:15, <213>
ORGANISM:Artificial Sequence
L:900 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:15,Line#:900
L:907 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:909 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:16, <213>
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ORGANISM:Artificial Sequence
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L:916 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
L:918 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:17, <213>
ORGANISM:Artificial Sequence
L:918 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:17, <213>
ORGANISM:Artificial Sequence
L:918 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:17,Line#:918
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